

Diffusion Maps

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Diffusion Maps



Diffusion Maps



Method Non-linear Spectral Dimensionality Reduction method.

Goal To describe properly the geometry of the data, i.e. to find a good distance in the original sample space that characterized well the relation between the points.

New data representation that will capture the main structure of the data in a few dimensions while preserving the local structure of the original data, i.e, reducing the sample distance to a Euclidean distance.

Main assumption The manifold metric can be approximated by the **diffusion distance** of a **Markov process** whose transition matrix is defined by an adequate normalization of the similarity matrix.



Continuous-Discrete Dictionary

Continuous version		Discrete version	
\mathcal{P}	operator	\mathbf{P}	matrix
f	function	\mathbf{v}	vector
$\phi_\ell(\mathbf{x}^{(i)})$	eigenfunction	$(\phi_\ell)_i$	eigenvector
\int	integral	\sum	summation



Motivation: Diffusion Processes



Diffusion Processes (I)



Diffusion (physical context)

The process by which a gas moves from regions of high density to regions of lower density according to the relative pressure of each region.

The diffusion equation is usually written:

$$\frac{\partial \chi}{\partial t} = c \nabla^2 \chi.$$

Diffusion (graph context)

The diffusion will be a model of spread across the graph.

Examples:

- The spread of an idea in a social network.
- The spread of a disease across some region.



Diffusion Processes (II)



Assumption Undirected weighted graph, with w_{ij} the similarity between vertex i and j .

The degree of each vertex is $d_i = \sum_j w_{ij}$.

Definition Let be χ_i a fluid or substance located in the nodes of the graph that flows from vertex j to an adjacent vertex i with a rate $c(\chi_j - \chi_i)$.

Diffusion constant: c .



Diffusion Processes (III)- Gas Diffusion Equation

- Flow in a short period of time: $c(\chi_j - \chi_i)dt$
- χ_i changes in a ratio:

$$\begin{aligned}
 \frac{\partial \chi_i}{\partial t} &= c \sum_j w_{ij}(\chi_j - \chi_i) \\
 &= c \sum_j w_{ij} \chi_j - c \chi_i \sum_j w_{ij} \\
 &= c \sum_j w_{ij} \chi_j - c \chi_i d_i \\
 &= c \sum_j (w_{ij} - \delta_{ij} d_i) \chi_j.
 \end{aligned}$$

- Matrix notation: $\frac{d\chi}{dt} = c(\mathbf{W} - \mathbf{D})\chi$.



Diffusion Processes (IV)- Laplacian Graph



Laplacian matrix

$$\mathbf{L} = \mathbf{D} - \mathbf{W}$$

We can then rewrite:

$$\frac{\partial \chi}{\partial t} + c\mathbf{L}\chi = 0.$$

We have arrived to the gas diffusion process equation (assuming $\mathbf{L} \approx \Delta = \nabla^2$).



Diffusion Processes (V)- Equation Solution

- Eigendecomposition of \mathbf{L} :

$$\mathbf{L}\phi_i = \lambda_i\phi_i.$$

- $\{\phi_i\}$ form an orthonormal basis, so:

$$\chi(t) = \sum_i w_i(t)\phi_i,$$

being $w_i(t)$ the coefficients to define χ depending on time t .

- Replacing in the diffusion equation:

$$\frac{\partial(\sum_i w_i(t)\phi_i)}{\partial t} + c\mathbf{L} \sum_i w_i(t)\phi_i = \sum_i \left(\frac{\partial w_i(t)}{\partial t} + c\lambda_i w_i(t) \right) \phi_i = 0.$$



Diffusion Processes (VI)- Equation Solution



Rewritten equation $\frac{\partial w_i(t)}{\partial t} + c\lambda_i w_i(t) = 0$

Solution $w_i(t) = w_i(0)e^{c\lambda_i t}$.

The flux can be determined under some initial conditions $w_i(0)$, just computing the eigendecomposition of **L**.

Diffusion Maps is a particular type of diffusion process that can be used to study the underlying relationship between points in a data set.



Diffusion Maps Algorithm



Defining Diffusion Coordinates: Building a Graph

- Starting sample: $\mathcal{S} = \{\mathbf{x}^{(1)}, \dots, \mathbf{x}^{(N)}\}$.
- A diffusion process over the data \Rightarrow **Random Walk** over a graph.
- First step: to build a **symmetric weighted graph**.

Affinity matrix

In terms of a kernel matrix \mathbf{K} which results from a kernel operator $\mathcal{K} : \mathcal{S} \times \mathcal{S} \rightarrow \mathbb{R}$, defining the matrix entries as $k_{ij} = \mathcal{K}(\mathbf{x}^{(i)}, \mathbf{x}^{(j)})$.

Properties:

- Symmetric: $k_{ij} = k_{ji}$.
- Positive: $k_{ij} \geq 0$.
- Positive semi-definite: $\mathbf{x}^{(i)\top} \mathbf{K} \mathbf{x}^{(i)} \geq 0 \quad \forall \mathbf{x}^{(i)} \in \mathcal{S}$.
- How to choose the kernel? It depends on the concrete problem.
 \Rightarrow Usual choice: **Gaussian kernel** defined as $k_{ij} = e^{\frac{-\|\mathbf{x}^{(i)} - \mathbf{x}^{(j)}\|^2}{2\sigma^2}}$.



Notebook

Building a Graph



Random Walk over the graph (I)

- We need to build now a **Markov Chain** (MC) over the graph.
- In particular, we build a **random walk**.
 - It defines a local relationship between points in the graph providing a structure to the data.

Normalized Graph Laplacian construction

We define a **transition probability** from the affinity matrix.

- 1 Degree of each node in the graph as:

$$d_i = \sum_{j \in \mathcal{S}} k_{ij}.$$

- 2 Transition probability \equiv normalized kernel:

$$p_{ij} = \frac{k_{ij}}{d_i}.$$



Random Walk over the graph (II)

Is this transition probability of the random walk well-defined?

① $\mathbf{P} \geq 0$

② $\sum_j p_{ij} = \frac{\sum_j k_{ij}}{d_i} = \frac{d_i}{d_i} = 1.$

- $\mathbf{P} \equiv$ probability of arriving from i to j in one step.
- $\mathbf{P}^T = (p_{ij}^T) \equiv$ probability of arriving from i to j in T steps.
- T : scale.
 - Running the process far away in time (with a higher T) lets us integrate the local geometry, so the structure of the data is revealed at different scales.



Notebook

Random Walk over the graph: Transition Probability and steps



Random Walk over the graph (III)

Stationary Distribution of the MC:

$$\pi_i = \frac{d_i}{\sum_k d_k}$$

Properties of π_i

- 1 We can prove that it is **stationary**:

$$\sum_i \pi_i p_{ij} = \sum_i \frac{d_i}{\sum_k d_k} \frac{k_{ij}}{d_i} = \sum_i \frac{k_{ij}}{\sum_k d_k} = \frac{d_j}{\sum_k d_k} = \pi_j.$$

- 2 It is a **reversible** chain, as it satisfies the balance equations:

$$\pi_i p_{ij} = \frac{d_i}{\sum_k d_k} \frac{k_{ij}}{d_i} = \frac{k_{ij}}{\sum_k d_k} = \frac{k_{ji}}{\sum_k d_k} = \frac{d_j}{\sum_k d_k} \frac{k_{ji}}{d_j} = \pi_j p_{ji}.$$

- 3 The chain is **irreducible**, as the graph is connected.
- 4 The chain is **aperiodic**: $g(\mathbf{x}^{(i)}) = \gcd\{r \geq 1 : p_{ii}^r > 0\} = 1 \forall i.$

\Rightarrow The MC is **ergodic**.

Notebook

Random Walk over the graph: MC Distribution π



Diffusion Distance (I)

Final objective To find a good distance in the original sample space that is reduced to a Euclidean distance.

Possible distance Based on the previous Markov chain.

$$\begin{aligned}
 \boxed{\mathcal{D}_T(\mathbf{x}^{(i)}, \mathbf{x}^{(j)})} &= \|p_{i\cdot}^T - p_{j\cdot}^T\|_{(1/\pi)}^2 = \sum_k \frac{(p_{ik}^T - p_{jk}^T)^2}{\pi_k} \\
 &= \sum_k \frac{(p_{ik}^T)^2 + (p_{jk}^T)^2 - 2p_{ik}^T p_{jk}^T}{\pi_k} = \sum_k \frac{p_{ik}^T p_{ik}^T + p_{jk}^T p_{jk}^T - p_{ik}^T p_{jk}^T - p_{jk}^T p_{ik}^T}{\pi_k} \\
 &= \sum_k \frac{1}{\pi_k} \left(p_{ik}^T p_{ki}^T \frac{\pi_k}{\pi_i} + p_{jk}^T p_{kj}^T \frac{\pi_k}{\pi_j} - p_{ik}^T p_{kj}^T \frac{\pi_k}{\pi_j} - p_{jk}^T p_{ki}^T \frac{\pi_k}{\pi_i} \right) \\
 &= \frac{1}{\pi_i} \sum_k p_{ik}^T p_{ki}^T + \frac{1}{\pi_j} \sum_k p_{jk}^T p_{kj}^T - \frac{1}{\pi_j} \sum_k p_{ik}^T p_{kj}^T - \frac{1}{\pi_i} \sum_k p_{jk}^T p_{ki}^T \\
 &= \frac{p_{ii}^{2T} - p_{ji}^{2T}}{\pi_i} + \frac{p_{jj}^{2T} - p_{ij}^{2T}}{\pi_j}.
 \end{aligned}$$



Diffusion Distance (II)

We will call this distance **Diffusion Distance**.

Properties of the Diffusion Distance

- ① $\mathcal{D}_T(\mathbf{x}^{(i)}, \mathbf{x}^{(j)})$ is symmetric.
 - ② $\mathcal{D}_T(\mathbf{x}^{(i)}, \mathbf{x}^{(j)})$ satisfies the triangle inequality.
- ⇒ It is a semi-metric.
- ⇒ If $k_{ij} > 0 \forall i, j$, it is a metric.
- ③ It is robust to noise (calculated as an average of all the paths of length t).

The **diffusion distance** measures the connectivity between two points in the dataset after $2t$ steps.

Intuition: $\mathcal{D}_T(\mathbf{x}^{(i)}, \mathbf{x}^{(j)})$ small if there exist a lot of paths that connect $\mathbf{x}^{(i)}$ and $\mathbf{x}^{(j)}$

⇒ when p_{ij}^{2t} and p_{ji}^{2t} are high.



Spectral Theory (I)



Theorem (Spectral Theorem)

Any symmetric matrix $\mathbf{A} \in \mathbb{R}^n \times \mathbb{R}^n$ can be reduced to an orthonormal basis determined by a diagonal matrix, i.e., $\mathbf{A} = \mathbf{\Phi} \mathbf{\Lambda} \mathbf{\Phi}'$, where

- $\mathbf{\Lambda} = \text{diag}\{\lambda_0, \lambda_1, \dots, \lambda_N\}$, with λ_ℓ the eigenvalues of \mathbf{A} ,
- $\mathbf{\Phi} = (\phi_0, \dots, \phi_N)$, with ϕ_ℓ the eigenvectors of \mathbf{A} .

! Our Markov matrix \mathbf{P} is not symmetric.

→ Let's consider its conjugated matrix \mathbf{A} :

$$a_{ij} = \frac{\sqrt{\pi_i}}{\sqrt{\pi_j}} p_{ij} = \frac{\sqrt{\pi_i}}{\sqrt{\pi_j}} \frac{k_{ij}}{d_i} = \frac{\sqrt{\frac{d_i}{\sum d_k}}}{\sqrt{\frac{d_j}{\sum d_k}}} \frac{k_{ij}}{d_i} = \frac{k_{ij}}{\sqrt{d_i} \sqrt{d_j}}$$



Spectral Theory (II)

- Applying the Spectral Theorem over \mathbf{A} :

$$a_{ij} = \sum_{\ell \geq 0} \lambda_{\ell} (\phi_{\ell})_i (\phi_{\ell})_j$$

- $\{\lambda_{\ell}\}_{\ell \geq 0}$, eigenvalues of \mathbf{A}
 - $\{\phi_{\ell}\}_{\ell \geq 0}$, eigenvector of \mathbf{A} .
- $\lambda_0 = 1$ corresponds to the eigenvector $\phi_0 = \sqrt{\pi}$.

$$(\mathbf{A}\sqrt{\pi})_i = \sum_j a_{ij} \sqrt{\pi_j} = \sum_j \frac{\sqrt{\pi_i}}{\sqrt{\pi_j}} p_{ij} \sqrt{\pi_j} = \sqrt{\pi_i} \sum_j p_{ij} = \sqrt{\pi_i} \mathbf{1} = \sqrt{\pi_i} \quad \forall i.$$



Spectral Theory (III)

- **P** rewritten:

$$p_{ij} = \frac{\sqrt{\pi_j}}{\sqrt{\pi_i}} a(\mathbf{x}^{(i)}, \mathbf{x}^{(j)}) = \frac{\sqrt{\pi_j}}{\sqrt{\pi_i}} \sum_{\ell \geq 0} \lambda_\ell (\phi_\ell)_i (\phi_\ell)_j = \sum_{\ell \geq 0} \lambda_\ell \frac{(\phi_\ell)_i}{\sqrt{\pi_i}} (\phi_\ell)_j \sqrt{\pi_j}.$$

⇒ Eigendecomposition of **P** for any r -steps:

$$p_{ij} = \sum_{\ell \geq 0} \lambda_\ell^r (\psi_\ell)_i (\varphi_\ell)_j,$$

where $(\psi_\ell)_i = \frac{(\phi_\ell)_i}{\sqrt{\pi_i}}$ and $(\varphi_\ell)_j = (\phi_\ell)_j \sqrt{\pi_j}$.

- $\lambda_0 = 1$ discarded also for **P**.
- The other eigenvalues will satisfy $|\lambda_\ell| < 1 \ \forall \ell \geq 1$.



Simplifying $\mathcal{D}_T(\mathbf{x}^{(i)}, \mathbf{x}^{(j)})$

$$\begin{aligned}
\boxed{\mathcal{D}_T(\mathbf{x}^{(i)}, \mathbf{x}^{(j)})} &= \|p_{ik}^T - p_{jk}^T\|_{(1/\pi)}^2 = \sum_k \frac{(p_{ik}^T - p_{jk}^T)^2}{\pi_k} \\
&= \sum_k \frac{\left(\sum_{\ell \geq 0} \lambda_\ell^T (\psi_\ell)_i (\varphi_\ell)_k - \sum_{\ell \geq 0} \lambda_\ell^T (\psi_\ell)_j (\varphi_\ell)_k\right)^2}{\pi_k} \\
&= \sum_k \frac{\sum_{\ell \geq 0} \lambda_\ell^{2T} ((\psi_\ell)_i - (\psi_\ell)_j)^2 (\varphi_\ell)_k^2}{\pi_k} = \sum_{\ell \geq 0} \lambda_\ell^{2T} ((\psi_\ell)_i - (\psi_\ell)_j)^2 \sum_k \frac{(\varphi_\ell)_k^2}{\pi_k} \\
&= \sum_{\ell \geq 0} \lambda_\ell^{2T} ((\psi_\ell)_i - (\psi_\ell)_j)^2 \sum_k \frac{((\varphi_\ell)_k \sqrt{\pi_k})^2}{\pi_k} \\
&= \boxed{\sum_{\ell \geq 0} \lambda_\ell^{2T} ((\psi_\ell)_i - (\psi_\ell)_j)^2}.
\end{aligned}$$



Diffusion Coordinates

- **Dimensionality Reduction:** $\lambda_\ell \rightarrow 0$ when ℓ grows (small contribution to the Diffusion Distance).
 - Given a precision δ , we work with:

$$\bar{M} = s(\delta, r) = \max\{\ell \in \mathbb{N} \text{ s.t. } |\lambda_\ell|^r > \delta |\lambda_1|^r\}.$$

- $\ell = 0$ is omitted as ψ_0 is constant.
- **Diffusion distance approximation:**

$$\mathcal{D}_r(\mathbf{x}^{(i)}, \mathbf{x}^{(j)}) \approx \sum_{\ell \geq 1}^{\bar{M}} \lambda_\ell^{2r} ((\psi_\ell)_i - (\psi_\ell)_j)^2.$$

Diffusion Coordinates

They are the natural coordinates for representing the original sample, and they are defined as:

$$\Psi_r(\mathbf{x}) = \begin{pmatrix} \lambda_1^r \psi_1(\mathbf{x}) \\ \vdots \\ \lambda_{\bar{M}}^r \psi_{\bar{M}}(\mathbf{x}) \end{pmatrix}.$$

Diffusion Maps

We call **Diffusion Map** to the family $\{\Psi_t\}_{t \in \mathbb{N}}$.

- These maps embed the data into the Euclidean space $\mathbb{R}^{\bar{M}}$:

$$\begin{aligned} \|\Psi_t(\mathbf{x}^{(i)}) - \Psi_t(\mathbf{x}^{(j)})\| &= \sum_{\ell \geq 1}^{\bar{M}} \lambda_\ell^{2t} ((\psi_\ell)_i - (\psi_\ell)_j)^2 \\ &\approx \mathcal{D}_t(\mathbf{x}^{(i)}, \mathbf{x}^{(j)}) = \sum_{\ell \geq 1}^{\bar{M}} \lambda_\ell^{2t} ((\psi_\ell)_i - (\psi_\ell)_j)^2 + \sum_{\ell > \bar{M}}^N \lambda_\ell^{2t} ((\psi_\ell)_i - (\psi_\ell)_j)^2. \end{aligned}$$



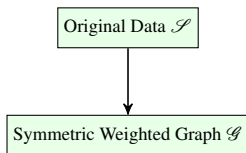
Diffusion Maps Algorithm Summary



Original Data \mathcal{S}



Diffusion Maps Algorithm Summary

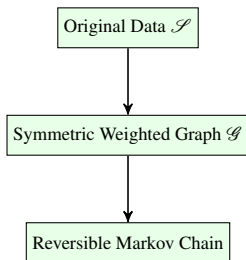


- Kernel definition:

$$\mathbf{K} = \{k_{ij}\} = \left\{ e^{\frac{-\|\mathbf{x}^{(i)} - \mathbf{x}^{(j)}\|^2}{2\sigma^2}} \right\}.$$



Diffusion Maps Algorithm Summary

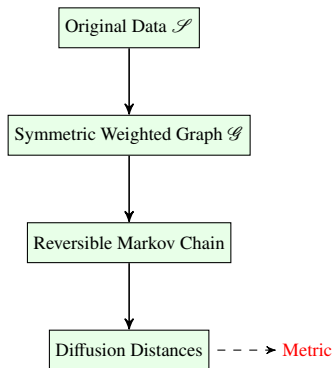


- Transition Probability Matrix definition:

$$p_{ij} = \left(\frac{k_{ij}}{d_i} \right) = \left(\frac{k_{ij}}{\sum_j k_{ij}} \right).$$



Diffusion Maps Algorithm Summary

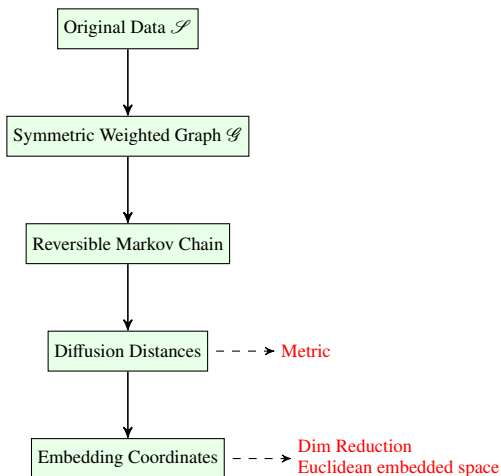


- Diffusion Distance:

$$\begin{aligned} \mathcal{D}_t(\mathbf{x}^{(i)}, \mathbf{x}^{(j)}) &= \|p_{i\cdot}^t - p_{j\cdot}^t\|_{(1/\pi)}^2 \\ &= \frac{p_{ii}^{2t} - p_{ji}^{2t}}{\pi_i} + \frac{p_{jj}^{2t} - p_{ij}^{2t}}{\pi_j}. \end{aligned}$$



Diffusion Maps Algorithm Summary



- Spectral Theory:

$$p_{ij}^T = \sum_{\ell \geq 0} \lambda_{\ell}^T (\psi_{\ell})_i (\varphi_{\ell})_j$$

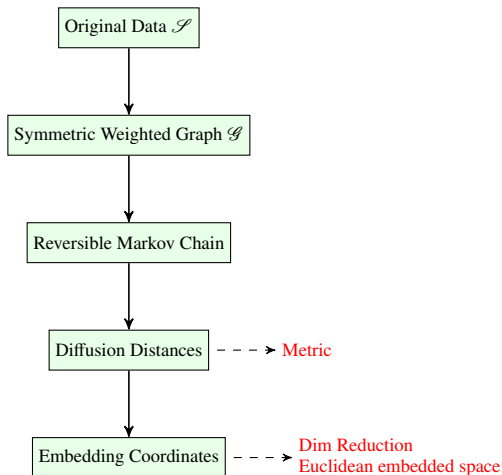
$$\rightarrow \mathcal{D}_T(\mathbf{x}^{(i)}, \mathbf{x}^{(j)}) = \sum_{\ell=0}^M \lambda_{\ell}^{2T} ((\psi_{\ell})_i - (\psi_{\ell})_j)^2.$$

- Dimensionality Reduction:

$$\bar{M} = \max\{\ell \in \mathbb{N} \text{ s.t. } |\lambda_{\ell}|^T > \delta |\lambda_1|^T\}.$$



Diffusion Maps Algorithm Summary



- Diffusion Coordinates Definition:

$$\Psi_t(\mathbf{x}) = \begin{pmatrix} \lambda_1^t \psi_1(\mathbf{x}) \\ \vdots \\ \lambda_M^t \psi_M(\mathbf{x}) \end{pmatrix}.$$

- Euclidean Distances over the embedding space:

$$\begin{aligned} \|\Psi_t(\mathbf{x}^{(i)}) - \Psi_t(\mathbf{x}^{(j)})\| &= \sum_{\ell=1}^{\bar{M}} \lambda_\ell^{2t} ((\psi_\ell)_i - (\psi_\ell)_j)^2 \\ &\approx \mathcal{D}_t(\mathbf{x}^{(i)}, \mathbf{x}^{(j)}). \end{aligned}$$



Density Influence



Density Influence



- The distribution of the sample data affects how the similarity matrix captures the local geometry of the data.
- The density is measured by the degree of the graph \mathbf{D} .
- We introduce a new parameter $\alpha \in [0, 1]$.
- Steps for normalizing:
 - 1 Define α -dependent density normalized matrix:

$$k_{ij}^{(\alpha)} = \frac{k_{ij}}{d_i^\alpha d_j^\alpha}.$$

- 2 Classical normalization:

$$p_{ij}^{(\alpha)} = \frac{k_{ij}^{(\alpha)}}{d_i^{(\alpha)}}.$$



Density Influence - Importance of α

$$\nabla_{\alpha} f = \frac{\Delta(f \mathbf{d}^{1-\alpha})}{\mathbf{d}^{1-\alpha}} - \frac{\Delta(\mathbf{d}^{1-\alpha})}{\mathbf{d}^{1-\alpha}} f$$

$$\alpha = 1 \quad \nabla_1 \equiv \Delta$$

→ DM captures the underlying geometry without interference from the sample's density \mathbf{d} .

$$\alpha = 0 \quad \nabla_0 f = \frac{\Delta(f \mathbf{d})}{\mathbf{d}} - \frac{\Delta(\mathbf{d})}{\mathbf{d}} f$$

→ \mathbf{d} influences how the diffusion coordinates capture the underlying geometry, unless \mathbf{d} is uniform ($\nabla_0 = \Delta$).



Tips on Practical Use



Tips on Practical Use (I)



- 1 The scale must be the same over all the features → Scale the data ($\mu = 0, \sigma = 1$).
- 2 Good hyper-parameter selection.

Affinity Matrix

- Gaussian kernel:

$$k_{ij} = e^{\frac{-\|\mathbf{x}^{(i)} - \mathbf{x}^{(j)}\|^2}{2\sigma^2}}.$$

- σ is crucial, it can be chose as a distance percentile (ρ):

$$\sigma = p_{\rho}(\|\mathbf{x}^{(i)} - \mathbf{x}^{(j)}\|)$$

- Usual ρ :
 - Median.
 - Maximum.



Tips on Practical Use (II)



Reduced dimension

For a given step τ :

$$\bar{M} = s(\delta, \tau) = \max\{\ell \in \mathbb{N} \text{ s.t. } |\lambda_\ell|^\tau > \delta |\lambda_1|^\tau\}.$$

Number of steps

By default, $\tau = 1$.

Density influence

By default, $\alpha = 1$.



Out-of-sample Extension



DM Advantages and Disadvantages



Advantages

- Powerful
- Elegant

Disadvantages

- Costly eigenanalysis of the transition matrix.
- The extension to new, unseen points (out-of-sample points) is not possible.



The Nyström Formula



- **Nyström**: approximate the eigenfunctions $\phi_j(\mathbf{x}^{(i)})$ of a symmetric and positive semidefinite kernel from the eigenvectors $(\phi_j)_i$ of a sample-based kernel matrix.
⇒ It maintains the eigenvalue and eigenvector convergence as the number of sample patterns grows.

- The Nyström Formula:

$$v_j(\mathbf{y}) = \frac{1}{\lambda_j} \sum_{i=1}^n v_j(\mathbf{x}^{(i)}) \mathcal{K}(\mathbf{y}, \mathbf{x}^{(i)})$$



The Nyström Formula applied to DM (I)

- Nyström approximates the DM embedding of new patterns
⇒ No need to repeat the eigendecomposition of \mathbf{P} over the training sample.
- Problem: \mathbf{P} is not symmetric.
⇒ Let's use its conjugate matrix \mathbf{A} .
- λ_j and $\phi_j(\mathbf{x}^{(i)})$: eigenvalues and eigenfunctions of \mathbf{A} of the sample data.
- Eigenvector extension for a new point \mathbf{x} :

$$\phi_j(\mathbf{x}) = \frac{1}{\lambda_j} \sum_{i=1}^N \phi_j(\mathbf{x}^{(i)}) a(\mathbf{x}, \mathbf{x}^{(i)}).$$



The Nyström Formula applied to DM (II)



Recall

- $a(\mathbf{x}, \mathbf{x}^{(i)}) = \frac{\pi(\mathbf{x})}{\pi(\mathbf{y})} \mathbf{P}(\mathbf{x}, \mathbf{x}^{(i)})$.
 - Eigenvalues: (λ_ℓ) .
 - Eigenvectors: (ϕ_ℓ) .
- \mathbf{P} left eigenvector: $(\psi_\ell)_i = \frac{(\phi_\ell)_i}{\sqrt{\pi_i}}$
- \mathbf{P} right eigenvector: $(\varphi_\ell)_j = (\phi_\ell)_j \sqrt{\pi_j}$



The Nyström Formula applied to DM (III)

- Nyström formula for a new \mathbf{x} yields:

$$\begin{aligned}\psi_j(\mathbf{x}) &= \frac{\phi_j(\mathbf{x})}{\sqrt{\pi(\mathbf{x})}} = \frac{1}{\sqrt{\pi(\mathbf{x})}} \left(\frac{1}{\lambda_j} \sum_{i=1}^N \phi_j(\mathbf{x}^{(i)}) a(\mathbf{x}, \mathbf{x}^{(i)}) \right) \\ &= \frac{1}{\lambda_j} \sum_{i=1}^N \phi_j(\mathbf{x}^{(i)}) \frac{a(\mathbf{x}, \mathbf{x}^{(i)})}{\sqrt{\pi(\mathbf{x})}} = \frac{1}{\lambda_j} \sum_{i=1}^N \phi_j(\mathbf{x}^{(i)}) \frac{\mathbf{P}(\mathbf{x}, \mathbf{x}^{(i)})}{\sqrt{\pi(\mathbf{x}^{(i)})}} \\ &= \frac{1}{\lambda_j} \sum_{i=1}^N \psi_j(\mathbf{x}^{(i)}) \mathbf{P}(\mathbf{x}, \mathbf{x}^{(i)}).\end{aligned}$$

⇒ The Nyström formula can be applied almost in its original form to \mathbf{P} .

- Moreover, this formulation is also valid for any α parameter value used to compute any of our $\mathbf{P}^{(\alpha)}$ matrices to be used in DM.



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